

Selected TF: AP1

TFName	AcceptLogOdds	AcceptProb	RejectCapability	AcceptCapability	BestMotifClusterId	CombineAvgScore	PositionBiasZscore	PeakRankBiasZscore	GlobalEnrichmentZscore
AP1	1.57429319091	0.828394775577	0.486	0.271	AP1_12	9.53281108867	11.2528443659	3.69771291138	9.06154725688
cluster: AP1_12	Enrichment Rank: 1.0	<p>69 CHIP-seq datasets</p> <p>Trained Model for Enrichment Rank</p>		<p>best motif: AP1_2</p>	<p>Position Distribution</p> <p>Peak-Rank Distribution</p> <p>Motif Position and Peak-Rank Distribution</p>				